

US-UY-866-60-SN

RESULT 2
US-08-899-514-1
Sequence 1, Application US/08899514
Patent No. 5910581
GENERAL INFORMATION:
APPLICANT: HABUCHI, OSAMI
APPLICANT: FUKUTA, MASAKAZU
TITLE OF INVENTION: POLYPEPTIDE OF GLYCOSAMINOGLYCAN
TITLE OF INVENTION: SULFOTRANSFERASE ORIGINATING FROM HUMAN AND DNA CODING
TITLE OF INVENTION: FOR THE POLYPEPTIDE
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: KNOBBE, MARTENS, OLSON & BEAR, LLP
STREET: 620 NEWPORT CENTER DRIVE, SIXTEENTH FLOOR
CITY: NEWPORT BEACH
STATE: CALIFORNIA
COUNTRY: US
ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/899,514
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:

FILING DATE:
 ATTORNEY/AGENT INFORMATION:
 NAME: DANIEL E ALTMAN
 REGISTRATION NUMBER: 34,115
 REFERENCE/DOCKET NUMBER: TOYAM21.001AUS
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 714 760 0404
 TELEFAX: 714 760 9502
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2156
 TYPE: nucleic acid
 STRANDEDNESS: both
 TOPOLOGY: linear
 MOLECULE TYPE: cDNA
 ORIGINAL SOURCE:
 ORGANISM: Human
 TISSUE TYPE: Fetal brain
 FEATURE:
 NAME/KEY: CDS
 LOCATION: 147..1583
 IDENTIFICATION METHOD: S

	Query Match	20.1%	Score 235.2:	DB 2,	Length 2156;
	Best Local Similarity	54.0%;	Pred. No. 7.4e-41;		
	Matches 589;	Conservative 0;	Mismatches 468;	Indels 33;	Gaps 4.
QY	91 gggccctatcccagccgagcgaggaatcgttgcacgttgctgtgtcctcgtg	150			
Db	510 GAGCCGCCACAGACCGGCGGTGGCGGGGCCCGGGCCACGTCGTCTCATGTGGCCACCAGC	569			
QY	151 cgcctcgggtcatccttccttttggccaaagtcttcaggccaaccgcgaacttcctactg	210			
Db	570 CGCACCGGCTCCTCGTGTGTGGCGCAAGTTCTTAACACGACGAGGCACACTTCTTACTTC	629			
QY	211 atggaagcccgagcgtgatgtgtggaaccaacctgtc-----gcaaggcagcg	258			
Db	630 TTTCGAGCCCGTGTTGGACATCGAGCCCAAGTGTCTTCGAGCCGGGGGGCCAACGCC	689			
QY	259 .gcaacgctcacatgtgccgtgagcgaactgtatgccttatcttttgtgcacatgac	318			
Db	690 GCGGGCTCGGCCCTGTGTATACGCGACGTGCTCAAGCAGCTCTTCTGTGGACCTGTAC	749			
QY	319 gtgtttgatgcctaca-----tgccacaagagcgaaacctgtccgccttttcac	369			
Db	750 GTGCTGGAGCACTTCATCAACGCCCGCTGCCCCGAGGACCACTGACTCAGTTCATGTTCGC	809			
QY	370 tgggcnaagcgccgagcgtgtgtctgcgcgcgcgcctgcagcgcccttccccagagcaac	429			
Db	810 CCGGGCTCCAGCCGCTCCTGTGTGCGAAGAACCCGCTGTACGCCCTTCGTCAAGAAGTTC	869			
QY	430 atcagcaagcagagacgtatgcagaagaactgtgcagcgcgacccattcagcctggcccg	489			
Db	870 TTCGAGAAGTACCACTGCACAGAACCCCGCTGCGGCCCTCAACGTGACGCTGGCCGA	929			
QY	490 gaggcctgcgctcctctaacagccaagtygtctcaaaggagtgcctcttcaacctgcag	549			
Db	930 GAGGCTCGCGCGCAAGSAGCACATTGGCCCTCAAGGCGGTGCCATCCGGCAGCTGGAG	989			
QY	550 gtgctctaacgctgtctcagcgaaaccgcgctcaaacctgcgcatcgtgcacctggtgcg	609			
Db	990 TTCCTGCAGCGCTGCGGAGGAGACCCCGCTGGAACCTGCCTCATTCAGCTGTGCCC	1049			
QY	610 gaccgcgagcgcgctgtcgtcctccgagagcgagcgagcccgatactgcacgcgacaac	669			
Db	1050 GACCCCGCGGCGTGTGCTGCGCTCGCCCATGTGTGSCCTTCGCCGCCAAGTAAAGACCTGG	1109			
QY	670 ggcatcgtctctgggcaccaaagtgtagtggtagggcggaacctcaaccttcgcttgat	729			
Db	1110 AAGAGTGGCTGACGACGAGGCGCCAGACGGCTTGAGGGAAGAGAGTGTACGCGGCTG	1169			

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RESULT 3
 US-08-655-878-1
 Sequence 1, Application US/08655878
 Patent No. 5827713
 GENERAL INFORMATION:
 APPLICANT: FUKUTA, MASAKAZU
 APPLICANT: HABUCHI, OSAMI
 TITLE OF INVENTION: DNA CODING FOR SULFOTRANSFERASE
 NUMBER OF SEQUENCES: 9
 CORRESPONDENCE ADDRESS:
 ADDRESSEE:
 STREET:
 CITY:
 STATE:
 COUNTRY:
 ZIP:
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/655,878
 FILING DATE:
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER:
 FILING DATE:
 ATTORNEY/AGENT INFORMATION:
 NAME:
 REGISTRATION NUMBER:
 REFERENCE/DOCKET NUMBER:
 TELECOMMUNICATION INFORMATION:
 TELEPHONE:
 TELEFAX:
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2354
 TYPE: nucleic acid
 STRANDEDNESS: both

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1      TOPOLOGY: 1linear
2      MOLECULE TYPE: cDNA to mRNA
3      ORIGINAL SOURCE:
4      ORGANISM: Chick
5      TISSUE TYPE: Embryo chondrocyte
6      FEATURE:
7      FEATURE: NAME/KEY: CDS
8      FEATURE: LOCATION: 211..1584
9      FEATURE: IDENTIFICATION METHOD: P
10     FEATURE:
11     FEATURE: NAME/KEY: sig_peptide
12     FEATURE: LOCATION: 211..309
13     FEATURE: IDENTIFICATION METHOD: P
14     FEATURE:
15     FEATURE: NAME/KEY: mat_peptide
16     FEATURE: LOCATION: 310..1584
17     FEATURE: IDENTIFICATION METHOD: P
18     FEATURE:
19     FEATURE: NAME/KEY: transmembrane domain
20     FEATURE: LOCATION: 280..321
21     FEATURE: IDENTIFICATION METHOD: P
22     FEATURE:
23     FEATURE: NAME/KEY: potential N-glycosylation site
24     FEATURE: LOCATION: 394..402
25     FEATURE: IDENTIFICATION METHOD: S
26     FEATURE:
27     FEATURE: NAME/KEY: potential N-glycosylation site
28     FEATURE: LOCATION: 427..435
29     FEATURE: IDENTIFICATION METHOD: S
30     FEATURE:
31     FEATURE: NAME/KEY: potential N-glycosylation site
32     FEATURE: LOCATION: 493..501
33     FEATURE: IDENTIFICATION METHOD: S
34     FEATURE:
35     FEATURE: NAME/KEY: potential N-glycosylation site
36     FEATURE: LOCATION: 916..924
37     FEATURE: IDENTIFICATION METHOD: S
38     FEATURE:
39     FEATURE: NAME/KEY: potential N-glycosylation site
40     FEATURE: LOCATION: 1405..1413
41     FEATURE: IDENTIFICATION METHOD: S
42     FEATURE:
43     FEATURE: NAME/KEY: potential N-glycosylation site
44     FEATURE: LOCATION: 1537..1545
45     FEATURE: IDENTIFICATION METHOD: S
46     US-08-655-878-1
47
48 Query Match 16.8%; Score 196.8; DB 1; Length 2354;
49 Best Local Similarity 52.4%; Pred. NO. 7.6e-33;
50 Matches 575; Conservative 0; Mismatches 487; Indels 36; Gaps 5;
51
52 QY 52 cagaccaccgcgcctcctcgtcttcatacatctcccgccaggccctcatccccagccgcgc 111
53   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
54 Db 475 CTGAGGACGCGGGCTGCGCAAGCTGACCTTGACATTGGCTGGGAGAGCTGGGCATAGCACCC 534
55
56 QY 112 ggcgagagatcgtgtgcacgctgctgtgtcctcgtgtgcgctcggtgcctcatcctctgt 171
57   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
58 Db 535 CCAGAGCCGCGGGCGGCATGCTCTGCTGATGGCCACACACGACGACCGGCTCTCTCTGCTT 594
59
60 QY 172 ggcacagctcttcagccacgaccccgacgctctctacactgatgagagcccgcttgcaagtgt 231
61   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
62 Db 595 GGGGAGTGTCTTCAACCCAGGCAACATATTTCACTCTTTGAGGCCCCCTGTGGCAGATC 654
63
64 QY 232 tggaccacacctgtc-----gcagggcaggcgccgaacgctgcacatgagccgtg 279
65   ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
66 Db 655 GAGAGGAGCGTCACTTTTGAGCCAGAGGGGGCCCAAGCGGCGGTGGCTCGGCCCTGCTGTAC 714
67
68 QY 280 cgcgaacctgatgcgctcatalcttttqtgagacatgagacggtttgagtcctacatgcaca 339
69   ||||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
70 Db 715 CGCGACGTGCTGCAGCAGCTCCTCTCTGCGACCTTACATTCTGTGAGAGCTTCACTCTCA 774
71
72 QY 340 cagagccggaacctgtcc-----gaccttttcaacttgggaaagaaacccacnnnnn 180

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15-00000-00000

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; RESULT 4
; Sequence 3, Application US/09263023
; Patent No. 6037159
; GENERAL INFORMATION:
; APPLICANT: Uchimura, Kenji
; APPLICANT: Muramatsu, Hideki
; APPLICANT: Kadomatsu, Kenji
; APPLICANT: Kannagi, Reiji
; APPLICANT: Habuchi, Osami
; APPLICANT: Muramatsu, Takashi
; TITLE OF INVENTION: POLYPEPTIDE OF N-ACETYLGALACTOSAMINE-6-0-SULFOTRANSFERASE AND
; FILE REFERENCE: TOYAMA41.001AUS
; CURRENT APPLICATION NUMBER: US/09/263,023
; CURRENT FILING DATE: 1999-03-05
; EARLIER APPLICATION NUMBER: JP 10-54007

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Query Match	12.7%;	Score 148.8;	DB 3;	Length 2409;
Best Local Similarity	50.7%;	Pred. No. 7.8e-23;		
Matches 563; Conservative	0;	Mismatches 457;	Indels 90;	Gaps 5.

[illegible]

Db 1506 caataactgtgtgtcggtacagagacactgtgtggagaccgccgtcaagacaactaagagaga 1565
QY 862 ctctaacgcttcaaccgagccctgaacctcaagccacaagctcgaggcctgtgattccacaacatc 921
Db 1566 gtgtacgatttgtgtggaactgtgtgagccccgaatgtagagcagtttgccttgaacatg 1625
QY 922 acccaaggtgtgggtatcggaagccaatgaggccttccataacttcgtctagggaatgag 981
Db 1626 accagtggtcg-----ggctcctcctcctcaagccttctgtgtatctctgacgcaatgac 1679
QY 982 cgaacgtctcccaagccctgagcccaagcgttgccttcaactaagatcctcgcgctgacag 1041
Db 1680 acgagggccgccaatgacctgagccgaccgctcacccttcagcagatcaacaagtgtag 1739
QY 1042 gaggtgtgagccgagcgagctgacagctgtgtgggtacaccgacctgtgtactctgcgacaa 1101
Db 1740 gagtttgcataccagcccaatgagccgtccctggtatgagcggtcaacagccctgagag 1799
QY 1102 cagcgtgaccccaaccctgtgactgtgtcgt 1131
Db 1800 gtcaagaccctcagcaagaccctgtctcg 1829

RESULT 5

US-09-263-023-1
Sequence 1, Application US/09263023
Patent No. 6037159

GENERAL INFORMATION:

APPLICANT: Uchimura, Kenji
APPLICANT: Muramatsu, Hideki
APPLICANT: Kadamatsu, Kenji
APPLICANT: Kanagaki, Reiji
APPLICANT: Habuchi, Osami
APPLICANT: Muramatsu, Takashi
TITLE OF INVENTION: POLYPEPTIDE OF N-ACETYLGLUCOSAMINE-6-0-SULFOTRANSFERASE AND
TITLE OF INVENTION: DNA ENCODING THE SAME
FILE REFERENCE: TOYAMA1.001AUS
CURRENT APPLICATION NUMBER: US/09/263, 023
EARLIER FILING DATE: 1999-03-05
EARLIER APPLICATION NUMBER: JP 10-54007
EARLIER FILING DATE: 1998-03-05
EARLIER APPLICATION NUMBER: JP 10-177844
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 1
LENGTH: 2150
TYPE: DNA
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: CDS
LOCATION: (470)...(1918)
US-09-263-023-1

Query Match 11.3%; Score 133; DB 3; Length 2150;
Best Local Similarity 54.3%; Pred. No. 1.5e-19;
Matches 324; Conservative 0; Mismatches 255; Indels 18; Gaps 2;

QY 109 ggcggcgagatcgtgtgcaagctgtgtgtgtctcctcgtgtgagctcggtcgtcattcctc 168
Db 806 ggaagggaagaagcgagctgt 865
QY 169 ttgggcaagctcttcaagcagcagcccgagcttctcaactgagtggagccgagtgagcat 228
Db 866 ttggtgagctcttcaacagaagcagtgagtggtcttctcctatagacctgtgtgagcac 925
QY 229 gtgtgacacacccgtgtcagagcgagcgcaacgctgcacatggccgtgtgcgacacctg 288
Db 926 gtgtgcaaaaactgttaccgccgggagagccggttccctgcaggggagcgagcgagacatg 985
QY 289 atgcgctctatcttctgtgcagacatgagcgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 338

Db 986 ctgagcgctctctaacgctgcgactcttgcgtttccagctgtatagccccgagcagcagt 1045
QY 339 -----acagagccgaacactgttccgaccttltcaactgaggcaacgagccgagctgtgc 393
Db 1046 gggggggcgcaaccctcaaccactctgggcatcttggggagcagccactaacaagtggtatgc 1105
QY 394 tcgcccgcgcctctgacgagccttcccccagagcaccatcagcaagcagagctatgcaag 453
Db 1106 tccctgcactctgtcctcctcctacagcaagaggtgtgtgactgtgtgagcagccgctg 1165
QY 454 ---aacctgtgacagcgagccactcagccttgcgagccgagagcctgcgctcctcagc 510
Db 1166 tgcataaagtgcccaactcacaacgcttgcacgcttcgagagagtgctgcgaagtacccgc 1225
QY 511 caagtgtgtcacaagaggtgtgccttcttcaacctgcaggtgtcttaccgctgtcagc 570
Db 1226 acggtgttatcaaggcggtgtccttctcgtatgtgtgtgtgtgtgtgtgtgtgtgtgt 1285
QY 571 gaccgcgcgtcaacctgcgacatcgtgacacctgtgtgcgagcccgagcgctgtgtgcgc 630
Db 1286 gatcagccttgcacctcaaggtcatcaccatgtagtgcctgtgtgtgtgtgtgtgtgtgt 1345
QY 631 tcccgggagagcgagcgagcccgatctggaagcgacaaagcagcatgtgtgtgtgtgtgt 687
Db 1346 tcccgcactccgcctgcgctcagcgctcatcctcggaagaccctacaggtgtgtgtgtgt 1402

RESULT 6

US-08-804-227C-1
Sequence 1, Application US/08804227C
Patent No. 5876991

GENERAL INFORMATION:

APPLICANT: DeHoff, Bradley S.
APPLICANT: Kuhstoss, Stuart A.
APPLICANT: Rostock, Paul R., Jr.
APPLICANT: Sutton, Kimberly L.
TITLE OF INVENTION: POLYPEPTIDE SYNTHASE GENES
CORRESPONDENCE ADDRESS: 15
ADDRESSEE: THOMAS G. PLANT 1501
STREET: LILLY CORPORATE CENTER
CITY: INDIANAPOLIS
STATE: IN
COUNTRY: USA
ZIP: 46285
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: ASCII(DOS) Text only
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/804,227C
FILING DATE: February 21, 1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Plant, Thomas, G.
REGISTRATION NUMBER: 35,784
REFERENCE/DOCKET NUMBER: X-8231
TELECOMMUNICATION INFORMATION:
TELEPHONE: 317-276-2459
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 43280 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
FEATURE:
NAME/KEY: CDS
LOCATION: 816..14234
FEATURE:
NAME/KEY: CDS
LOCATION: 14351..19945

Fri Feb 1 20:20:57 2002

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Page 6

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:      FEATURE:      CDS
:      NAME/KEY:      20010..31199
:      LOCATION:
:      FEATURE:
:      NAME/KEY:      CDS
:      LOCATION:      31232..36067
:      FEATURE:
:      NAME/KEY:      CDS
:      LOCATION:      36249..41774
:
US-08-804-227C-1

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Query Match	5.58;	Score 64.6;	DB 2;	Length 43280;
Best Local Similarity	44.78;	Pred. No. 4.4e-05;		
Matches 305; Conservative	0;	Mismatches 369;	Indels 9;	Gaps 1,

[illegible]

RESULT 7
 US-09-040-984-63
 Sequence 63, Application US/09040984
 Patent No. 6210883
 GENERAL INFORMATION:
 APPLICANT: Reed, Steven G.
 APPLICANT: Wang, TongTong
 TITLE OF INVENTION: COMPOUNDS AND

TITLE OF INVENTION: OF LUNG CANCER
 NUMBER OF SEQUENCES: 86
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: SEED AND BERRY LLP
 STREET: 6300 Columbia Center, 701 Fifth Avenue
 CITY: Seattle
 STATE: WA
 COUNTRY: USA
 ZIP: 98104
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette
 COMPUTER: IBM Compatible
 OPERATING SYSTEM: DOS
 SOFTWARE: FASTSEQ for Windows Version 2.0
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/040,984
 FILING DATE: 18-MAR-1998
 CLASSIFICATION:
 ATTORNEY/AGENT INFORMATION:
 NAME: Makl, David J.
 REGISTRATION NUMBER: 31,392
 REFERENCE/DOCKET NUMBER: 210121.456
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 206-622-4900
 TELEFAX: 206-282-6031
 TELEX:
 INFORMATION FOR SEQ ID NO: 63:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 731 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 TOPOLOGY: linear

Query Match	4.98;	Score 57.8;	DB 4;	Length 731;
Best Local Similarity	58.48;	Pred. No. 0.00061;		
Matches 101; Conservative	0;	Mismatches 72;	Indels 0;	Gaps 0;

[illegible]

RESULT 8
 US-08-125-468-1/c
 ; Sequence 1, Application US/08125468
 ; Patent No. 5589385
 ;
 ; GENERAL INFORMATION:
 ;
 APPLICANT: Ryan, Michael J.
 APPLICANT: Lotvin, Jason A.
 APPLICANT: Strathy, Nancy
 APPLICANT: Fantini, Susan E.
 TITLE OF INVENTION: Cloning of the biosynthetic pathway for
 TITLE OF INVENTION: chlorotetracycline and tetracycline formation and cosmi
 TITLE OF INVENTION: useful therein
 NUMBER OF SEQUENCES: 1
 ;
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: American Cyanamid Company
 STREET: One Cyanamid Plaza
 CITY: Wayne
 STATE: New Jersey
 COUNTRY: USA
 ZIP: 07470
 ;
 COMPUTER READABLE FORM:

SOFTWARE: FastSeq for Windows Version 3.0

SEQ ID NO 2
LENGTH: 1931
TYPE: DNA
ORGANISM: EBNA
US-09-130-114-2

Query Match
Best Local Similarity 44.7%; Score 55.6; DB 2: Length 1931;
Matches 217; Conservative 0; Mismatches 269; Indels 0; Gaps 0;

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QY 476 tcagctggccgagagcctgcgcctctctacagcagctgtgtctcaaggagtgct 535
    |||||
DB 450 tctctctcccgctctctcccgctctctctctctctctctctctctctcccgctcc 509

QY 536 tcttaacctgacagtgctctacacgctgtctacagcagcccgctcaacctgcgcatcg 595
    |||||
DB 510 tcttctctctcccgctctctctctctctctctctctctctctctctctctctcccg 569

QY 596 tgcacctgtgtgcgaccccgagcgctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 655
    |||||
DB 570 tctctcccgctctctctctctctctctctctctctctctctctctctctctctcccg 629

QY 656 tggcagcgacagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 715
    |||||
DB 630 tctctcccgctctctctctctctctctctctctctctctctctctctctctctcccg 689

QY 716 acctgagcctgattctgcgagtggtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 775
    |||||
DB 690 tctctgtctctcccgctctctctctctctctctctctctctctctctctctctcccg 749

QY 776 tcaagccgcacacctctctctctctctctctctctctctctctctctctctctctcc 835
    |||||
DB 750 tctctcccgctctctctctctctctctctctctctctctctctctctctctctcccg 809

QY 836 gggagcgctgtgcagagatctgcgacactctacacgctgtgtgtgtgtgtgtgtgtgt 895
    |||||
DB 810 tctctctctctcccgctctctctctctctctctctctctctctctctctctctcccg 869

QY 896 agctcagcgctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 955
    |||||
DB 870 tctctgtctctcccgctctctctctctctctctctctctctctctctctctctcccg 929

QY 956 cctctcc 961
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DB 930 cctctcc 935
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RESULT 11

US-09-105-537-34
Sequence 34, Application US/09105537A
Patent No. 6265202
GENERAL INFORMATION:
APPLICANT: Sherman, D.H.
APPLICANT: Liu, H.
APPLICANT: Xue, Y.
APPLICANT: Zhao, L.
TITLE OF INVENTION: DNA encoding methymycin and pikromycin
FILE REFERENCE: 600.438US1
CURRENT APPLICATION NUMBER: US/09/105,537A
CURRENT FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 34
LENGTH: 4689
TYPE: DNA
ORGANISM: Streptomyces venezuelae
US-09-105-537-34

Query Match
Best Local Similarity 44.7%; Score 54.6; DB 4: Length 4689;
Matches 225; Conservative 0; Mismatches 284; Indels 0; Gaps 0;

Matches 225; Conservative 0; Mismatches 284; Indels 0; Gaps 0;

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QY 582 caacctgcgcatcgtgcacctgtgtgcgaccccgagcgctgtgtgtgtgtgtgtgtgtgt 641
    |||||
DB 3408 ccaccacgagccgaacacacctctctctctctctctctctctctctctctctctct 3467

QY 642 ggcgggcccagatctgtgcagcgacacagcgatcgtgtgtgtgtgtgtgtgtgtgtgt 701
    |||||
DB 3468 caaccaactcacccgcgaactcacccgcatcgtgtgtgtgtgtgtgtgtgtgtgtgt 3527

QY 702 ggaagccgacacctacacctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 761
    |||||
DB 3528 cgtcgcgacccccacacgcatgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 3587

QY 762 cgagcgccacacactcacagccgacacacctctctctctctctctctctctctctct 821
    |||||
DB 3588 caccgcgctgtctcacacacccgacgacccgagcgatcgtgtgtgtgtgtgtgtgtgt 3647

QY 822 cgagacacctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 881
    |||||
DB 3648 ggaagacatcgcgccacatctctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 3707

QY 882 gacctcacgcacacagctgcagcgctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 941
    |||||
DB 3708 gctcgcgacacctcgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 3767

QY 942 caagcgaatcgagcgctctacatctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 1001
    |||||
DB 3768 cagcgacgacgagcgagcgctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 3827

QY 1002 ggcacagcgctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 1061
    |||||
DB 3828 ggcgcgagcccgagcgagcgagcgagcgagcgagcgagcgagcgagcgagcgagcg 3887

QY 1062 gcaagctgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgtgt 1090
    |||||
DB 3888 catggagcgagcgagcgagcgagcgagcgagcgagcgagcgagcgagcgagcgag 3916
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RESULT 12

US-09-105-537-5
Sequence 5, Application US/09105537A
Patent No. 6265202
GENERAL INFORMATION:
APPLICANT: Sherman, D.H.
APPLICANT: Liu, H.
APPLICANT: Xue, Y.
APPLICANT: Zhao, L.
TITLE OF INVENTION: DNA encoding methymycin and pikromycin
FILE REFERENCE: 600.438US1
CURRENT APPLICATION NUMBER: US/09/105,537A
CURRENT FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 43
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 5
LENGTH: 36778
TYPE: DNA
ORGANISM: Streptomyces venezuelae
US-09-105-537-5

Query Match
Best Local Similarity 44.7%; Score 54.6; DB 4: Length 36778;
Matches 225; Conservative 0; Mismatches 284; Indels 0; Gaps 0;

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QY 582 caacctgcgcatcgtgcacctgtgtgcgaccccgagcgctgtgtgtgtgtgtgtgtgtgt 641
    |||||
DB 30398 ccaccacgagcgagcgaacacacctctctctctctctctctctctctctctctctct 30457

QY 642 ggcgggcccagatctgtgcagcgacacagcgatcgtgtgtgtgtgtgtgtgtgtgtgt 701
    |||||
DB 30458 caccacactcacccgcgaactcacccgcatcgtgtgtgtgtgtgtgtgtgtgtgtgt 30517
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[illegible]

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RESULT 13
US-09-320-878-19
: Sequence 19, Application US/09320878A
: Patent No. 6117659
: GENERAL INFORMATION:
: APPLICANT: ASHLEY, Gary
: APPLICANT: BETLACH, Melanie C.
: APPLICANT: BETLACH, Mary C.
: APPLICANT: McDANIEL, Robert
: APPLICANT: TANG, Li
: TITLE OF INVENTION: RECOMBINANT NARBONOLIDE POLYKETIDE SYNTHASE
: FILE REFERENCE: 300622002120
: CURRENT APPLICATION NUMBER: US/09/320,878A
: CURRENT FILING DATE: 1999-05-27
: EARLIER APPLICATION NUMBER: CIP OF 09/141,908
: EARLIER FILING DATE: 1998-08-28
: EARLIER APPLICATION NUMBER: CIP OF 09/073,538
: EARLIER FILING DATE: 1998-05-06
: EARLIER APPLICATION NUMBER: CIP OF 08/846,247
: EARLIER FILING DATE: 1997-04-30
: EARLIER APPLICATION NUMBER: 60/119,139
: EARLIER FILING DATE: 1999-02-08
: EARLIER APPLICATION NUMBER: 60/100,880
: EARLIER FILING DATE: 1998-09-22
: EARLIER APPLICATION NUMBER: 60/087,080
: EARLIER FILING DATE: 1998-05-28
: NUMBER OF SEQ ID NOS: 34
: SOFTWARE: PatentIn Ver. 2.0
: SEQ ID NO 19
: LENGTH: 38506
: TYPE: DNA
: ORGANISM: Streptomyces venezuelae
: US-09-320-878-19

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Query Match	4.7%;	Score 54.6;	DB 3;	Length 38506;
Best Local Similarly	44.28;	Pred. NO. 0.0053;		
Matches 225; Conservative	0;	Mismatches 284;	Indels 0;	Gaps 0;

OY 582 caacctgcgcatcgtgcacactygtgcgcgaaccgcggccgtgtctgcgtcccggagagc 641
+ + + + +
Db 28540 ccaccacggaggcgaaacacctctctctcttcgtcagccgcagcgcggaacaagcccccgagc 28599

OY 642 ggcgcgccgatactgtgcacgcgcacaacgcgcatactgtctgtggcaaccaagcgcaagtggat 701

[illegible]

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US-09-050-863-2

Query Match 4.5%; Score 52.8; DB 3; Length 2580;
Best Local Similarity 42.9%; Pred. No. 0.0082;
Matches 316; Conservative 0; Mismatches 417; Indels 3; Gaps 1;

QY 400 cccgctgagcgccttcccgagagcaccatcagcaagcagatagcaagacactg 459
DB 1500 CTCCTCCCTGGCTCTTTCAGACTCCCTCCCTGCTCTTTCAGCTCTTACCCCGGC 1441
QY 460 tgcagcgcagcaccatcagcctgagcagcagcctgagcctcctacagcagctgtg 519
DB 1440 GGCTCCACTACCTCTCTGACCCCGGCTCCACTACTCTCTGACCCCGGCTCCACTGC 1381
QY 520 ctcaagagagtgctcttctcaactgagtgctctacccgctgctcagcagcccgcg 579
DB 1380 CTCCTCCAGCCCGGCTCCACTCTCTCTGCTCTCTGCTCTCTGCTCTCTCTCTG 1321
QY 580 ctcaactgagcctgctgacactgctgctgagcagcagcagcagcagcagcagcag 639
DB 1320 CTCCTCCCTCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTC 1261
QY 640 gcgagcgcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 699
DB 1260 CTGCCCCCTCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTCTCTGCTC 1201
QY 700 gtgagagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 759
DB 1200 CTCCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 1141
QY 760 gccgagcgcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 819
DB 1140 CCCCCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 1084
QY 820 ttcgagagcctgagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 879
DB 1083 CTCCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 1024
QY 880 ctgacccctcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 939
DB 1023 CTGCCCCCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 964
QY 940 ggcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 999
DB 963 CTGCCCCCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 904
QY 1000 tggcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 1059
DB 903 CTCCTCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTG 844
QY 1060 ctgagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 1119
DB 843 CCCCCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 784
QY 1120 gatctgt 1135
DB 783 CTCCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTCTGCTC 768

RESULT 15

US-09-130-114-1

Sequence 1, Application US/09130114

Patent No. 5976807

GENERAL INFORMATION:

APPLICANT: Horlick, Robert A.

APPLICANT: Damaj, Bassam B.

APPLICANT: Robbins, Alan K.

TITLE OF INVENTION: Eukaryotic Cells Stably Expressing Genes

FILE REFERENCE: 0867/1D903US1

CURRENT APPLICATION NUMBER: US/09/130,114

CURRENT FILING DATE: 1998-08-06

NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 1
LENGTH: 5452
TYPE: DNA
ORGANISM: YEBNA
US-09-130-114-1

Query Match 4.5%; Score 52.8; DB 2; Length 5452;
Best Local Similarity 42.9%; Pred. No. 0.0092;
Matches 316; Conservative 0; Mismatches 417; Indels 3; Gaps 1;

QY 400 cccgctgagcgccttcccgagagcaccatcagcaagcagatagcaagacactg 459
DB 1305 ctctcccttgctcttcaagacttcccccctgctcttcaagctctaccccgcg 1364
QY 460 tgcagcgcagcaccatcagcctgagcagcagcagcagcagcagcagcagcagcag 519
DB 1365 ggcctcactactcctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1424
QY 520 ctcaagagagtgctcttctcaactgagtgctctacccgctgctcagcagcccgcg 579
DB 1425 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1484
QY 580 ctcaactgagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 639
DB 1485 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1544
QY 640 gcgagcgcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 699
DB 1545 ctgcccctctgctctgctctgctctgctctgctctgctctgctctgctctgctctg 1604
QY 700 gtgagagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 759
DB 1605 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1664
QY 760 gccgagcgcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 819
DB 1665 cccctctgctctgctctgctctgctctgctctgctctgctctgctctgctctgctc 1721
QY 820 ttcgagagcctgagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 879
DB 1722 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1781
QY 880 ctgacccctcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 939
DB 1782 ctgcccctcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 1841
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QY 1000 tggcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagc 1059
DB 1902 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 1961
QY 1060 ctgagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 1119
DB 1962 cccctcctcagcagcagcagcagcagcagcagcagcagcagcagcagcagcagcag 2021
QY 1120 gatctgt 1135
DB 2022 ctctcagaccccgctcagcagcagcagcagcagcagcagcagcagcagcagcagc 2077

Search completed: January 31, 2002, 18:11:04
Job time: 10808 sec